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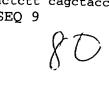
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     gaacaaatac aatggatgac aggcaaccca cctgtcccag tgggagaaat gtataaaaga 780
                                          SEQ 34
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    agggcccta gaaaaaaggg ctgttggaag tgtggaagag agggacacca aatgaaggac 1260
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    aggaaagcca tattaggaca taaaattagc cctaggtgta actatcaagc aggacataac 420
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    aaggtaggat ctctacaata cttggcacta acagcattaa tagctccaaa aaagacaaag 480
    ccgcctttgc ctagtgtcca gaaactagta gaagacagat ggaacaagcc ccagaagacc 540
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    <211> 291
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    ataaggatat tgcaacaact actgtttatc catttcagaa ttgggtgccg tcatagcaga 240
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    <211> 2610
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    <213> Human immunodeficiency virus type 1
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     agcagtcaga tacatcaaga ttttgtacca aagcagtaag tattgttaag catatgtaat 240
     gtcaaatttg ttagcaatag gcatagcagc attaatagta gcactaataa taacaatagt 300
     tgtgtggact atagcatata tagaatataa gaaactggta aggcaaagaa aaataaatag 360
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     agaggaattg gcagcacttg gggaagtggg gccttttatt cctggggaca ttaataatct 480
     gtaatgctgc agaaaactta tgggttacag tctattatgg ggtacctgtg tggaaagaag 540
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     aaagaatatc tttaggacca ggacgagtat tttatacaac aggagaaata ataggagaca 1380
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SEQ 36

107. D

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    atagaatata agaaactggt aaggcaaaga aaaataaata ggttatataa aagaataagc 240
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    ggggaagtgg ggccttttat tcctggggac attaataatc tgtaatgctg cagaaaactt 360
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     tgtacccaca gatcccaatc cacaagaagt agttctggaa aatgtaacag aaaggtttaa 540
     tatgtgggaa aataacatgg tagaacaaat gcatacagat ataatcagtt tatgggatca 600
     aagcctaaag ccatgtgtga agttaacccc actctgtgtt actttagatt gtagaaacat 660
     tgccaccaat ggcaccaatg acactattgc catcaatgac actctgaagg aagatccaga 720
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<212> DNA

<213> Human immunodeficiency virus type 1

SEO 37

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     aggttatata aaagaataag cgaaagagca gaagacagtg gcaatgagag tgagggggat 180
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     gtacctgtgt ggaaagaagc aaccactact ctattctgtg catcagatgc taaatcatat 180
     gaaaaagagg cacataatgt ctgggctaca catgcttgtg tacccacaga tcccaatcca 240
     caagaagtag ttctggaaaa tgtaacagaa aggtttaata tgtgggaaaa taacatggta 300
25
     gaacaaatgc atacagatat aatcagttta tgggatcaaa gcctaaagcc atgtgtgaag 360
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     actattgcca tcaatgacac tctgaaggaa gatccagagg caatacaaaa ctgttctttc 480
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     gatatagtac aaatcaacaa ggatgacaat agaacataca gactaataaa ttgtgatgcc 600
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     tcaaccatta cacaggcttg tccaaaggta tcttgggatc caattcccat acattattgt 660
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     aattggtttg acataacaca gtggctgtgg tatataaaaa tattcataat gatagtagga 2040
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                                         SEQ 38
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     ctggggagat ttgcacttaa ccgtgacctt ttagaaacag cagaaggttg tgtgcaaata 180
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SEQ 39

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SEQ 40

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     tatagaaatt aattgtatca gacccaataa caatacaaga aaaagtatac caatcggacc 1320
     aggacaagcg ttctatgcaa caggtgatat aataggagac ataagacaag cacattgtaa 1380
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SEQ 48

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SEO 50

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SEQ 57

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SEO 59

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SEQ 63

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SEO 65

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SEQ 75

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SEO 76

147

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SEQ 85

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156

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13

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J/

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SEQ 93

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